

CACACNTCCCCTACACATAGATATACATACAAAATCACAGAAAGTAATAGATATGGCTAAGTTTGCTACCAT

M A K F A T I

CATCTCTTCTCTTTGCTGCTCTTTGTTCTCTTTGCTGCCCTTTGAAGCACCAACAATGGTGGATGCAAGGTT

I S L L F A A L V L F A A F E A P T M V D A R L

Δ

GTGCGAGAGACCAAGTGGGACATGGTCAGGAGTTTGTGGGAACAACAATGCATGCAGGAACCAATGCAGAAA

C E R P S G T W S G V C G N N A C R N Q C R N

CCTTGAAAGAGCAGAACACGGATCTTGCAACTATGTCTTCCAGCTCACAAAATGTATTGTACTTCCCATG

L E R A E H G S C N Y V F P A H K C I C Y F P C

TTAATCTACCAAATCACTTTTGTGCTTGTGTGTATTTTACATGTTATGTGTTTATTACATGAAATAAG

\*

TCTGTGTCATCCTTATGGGTGACCTTATGACATGTACCAGATATATCATATATGTATGTTGTTGTGTGT

GGCAATTATAAACTTTTATTTGTGGATGCAAAAAAAAAAAAAAAAAAAA

FIG. 1

|         |  |
|---------|--|
| Al yAFP | <u>MAKFATIIISLLFAALVLF</u> <u>AAFEAPT</u> <u>MDA</u> -RLCERPSGTWSGVCNNNACRNQC                  |
| Rs-AFP1 | <u>MAKFASIIALLFAALVLF</u> <u>AAFEAET</u> <u>VEA</u> <u>QKLCERPSGTWSGVCNNNACKNQC</u>            |
| Rs-AFP2 | <u>MAKFASIIALLFAALVLF</u> <u>AAFEAET</u> <u>VEA</u> <u>QKLCQRP</u> SGTWSGVCNNNACKNQC<br>** * * |

|         |                  |                       |
|---------|------------------|-----------------------|
| Al yAFP | RNLEAEHGSCNYVFP  | PAHKCICYFPC           |
| Rs-AFP1 | INLEKARHGSCNYVFP | PAHKCICYFPC           |
| Rs-AFP2 | IRLEKARHGSCNYVFP | PAHKCICYFPC<br>** * * |

FIG. 2

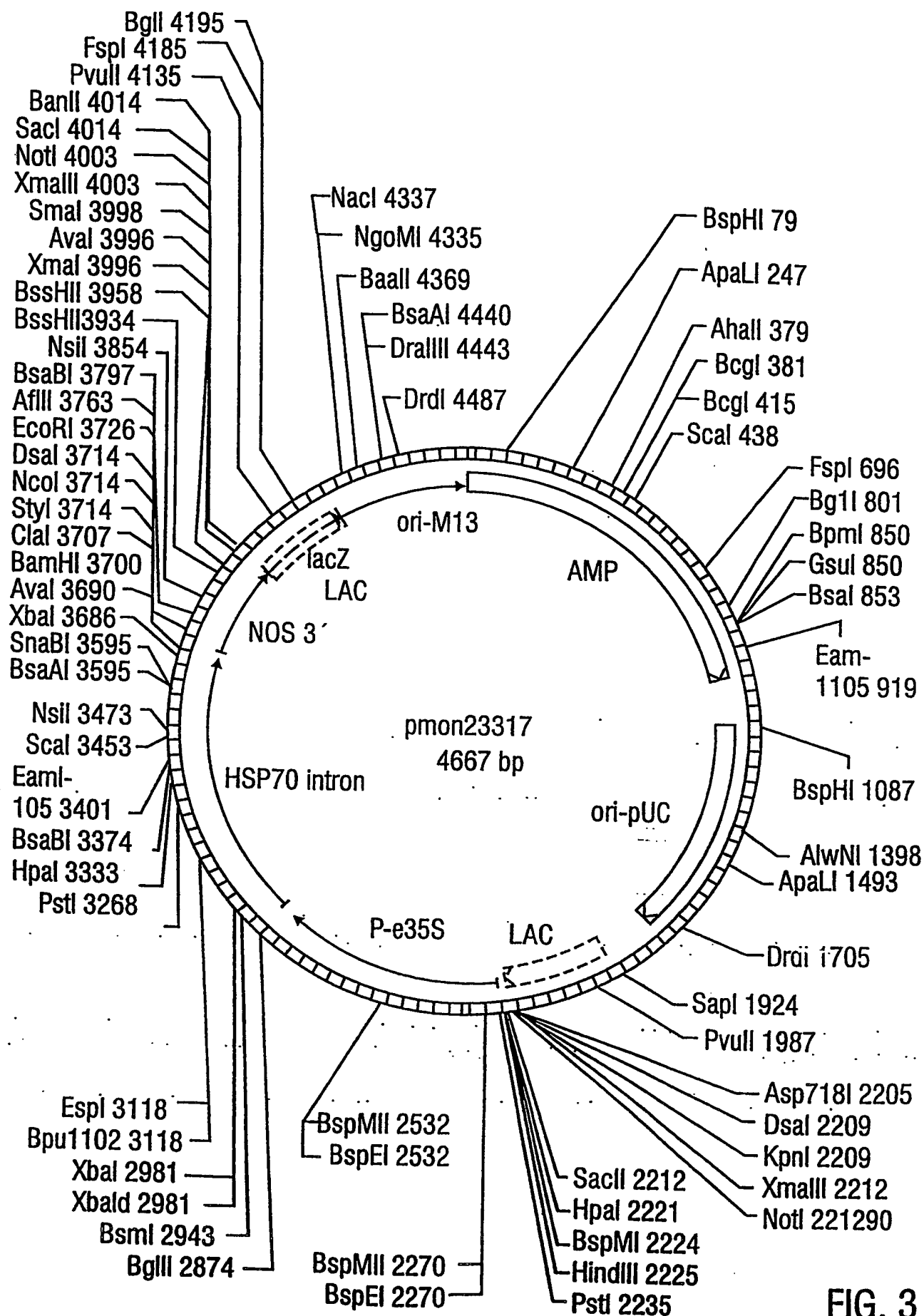


FIG. 3

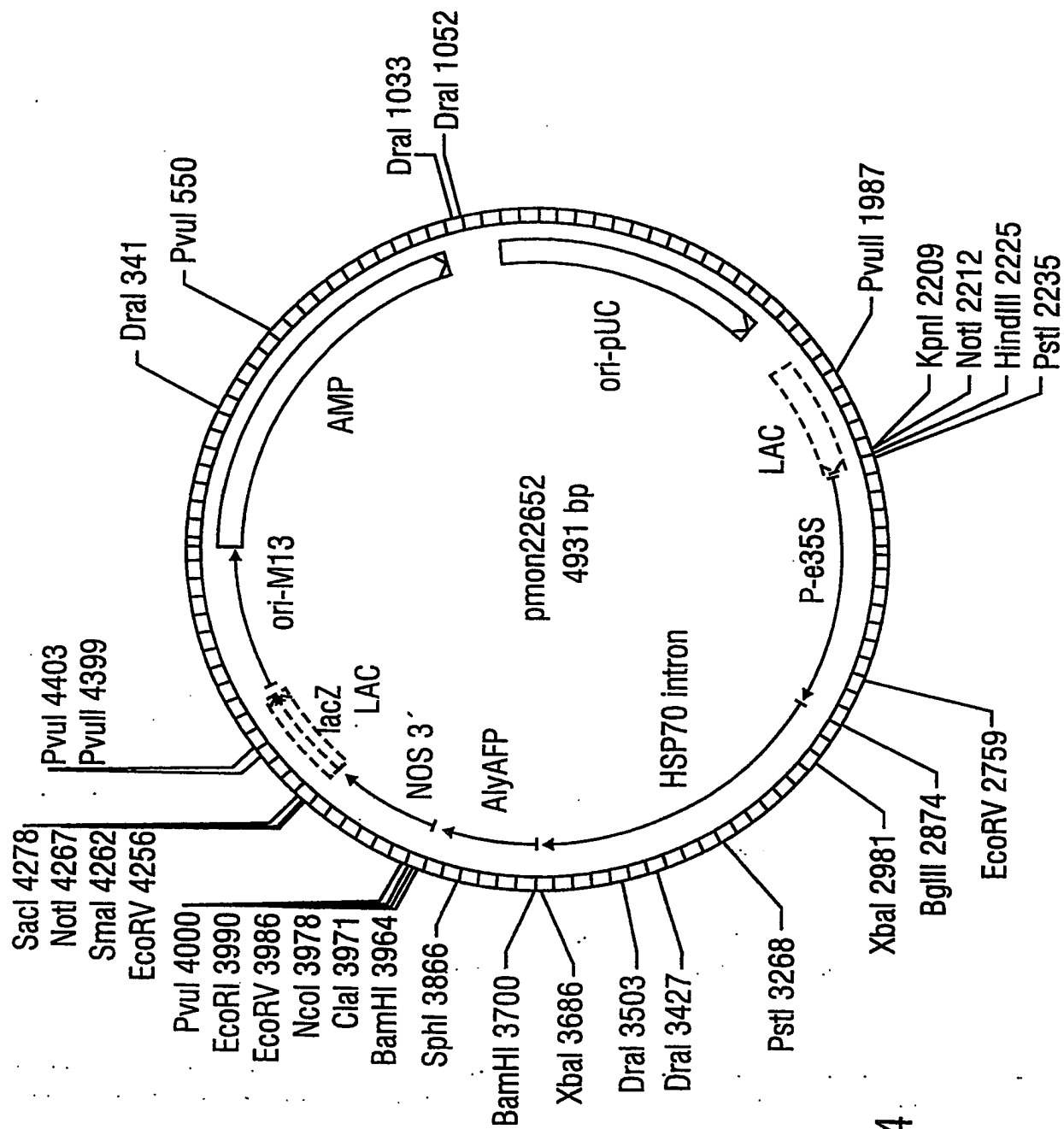


FIG. 4

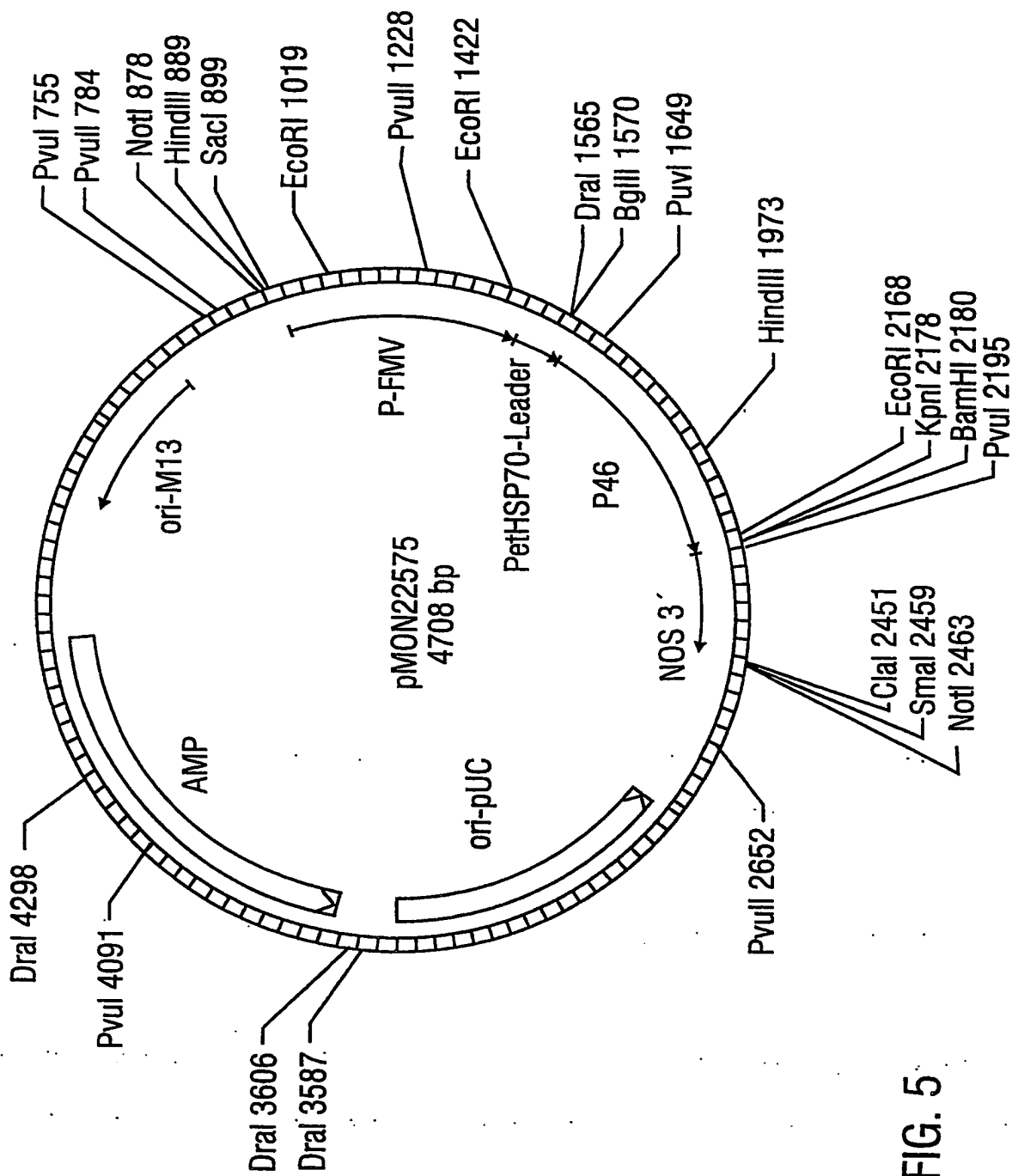


FIG. 5

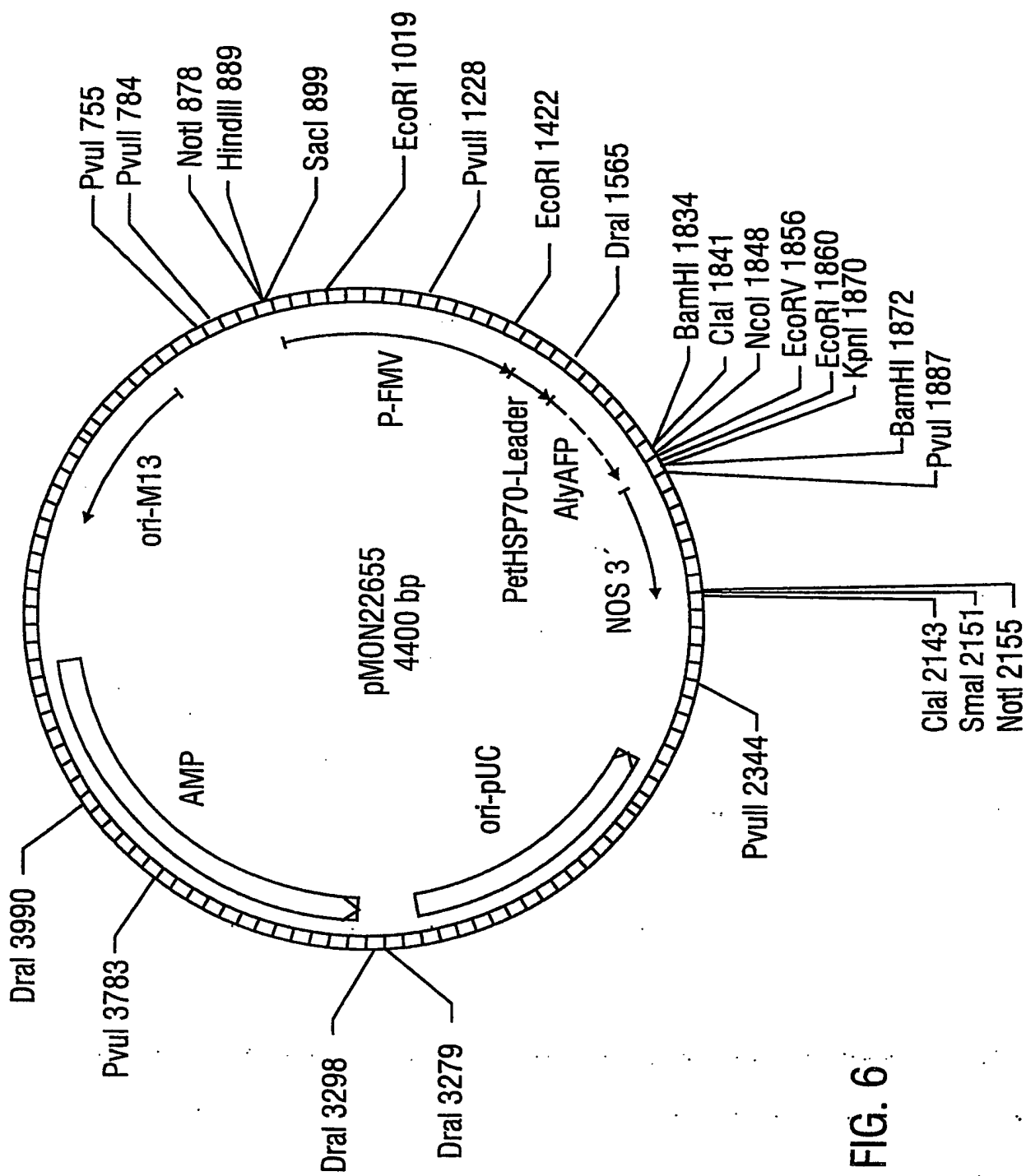


FIG. 6

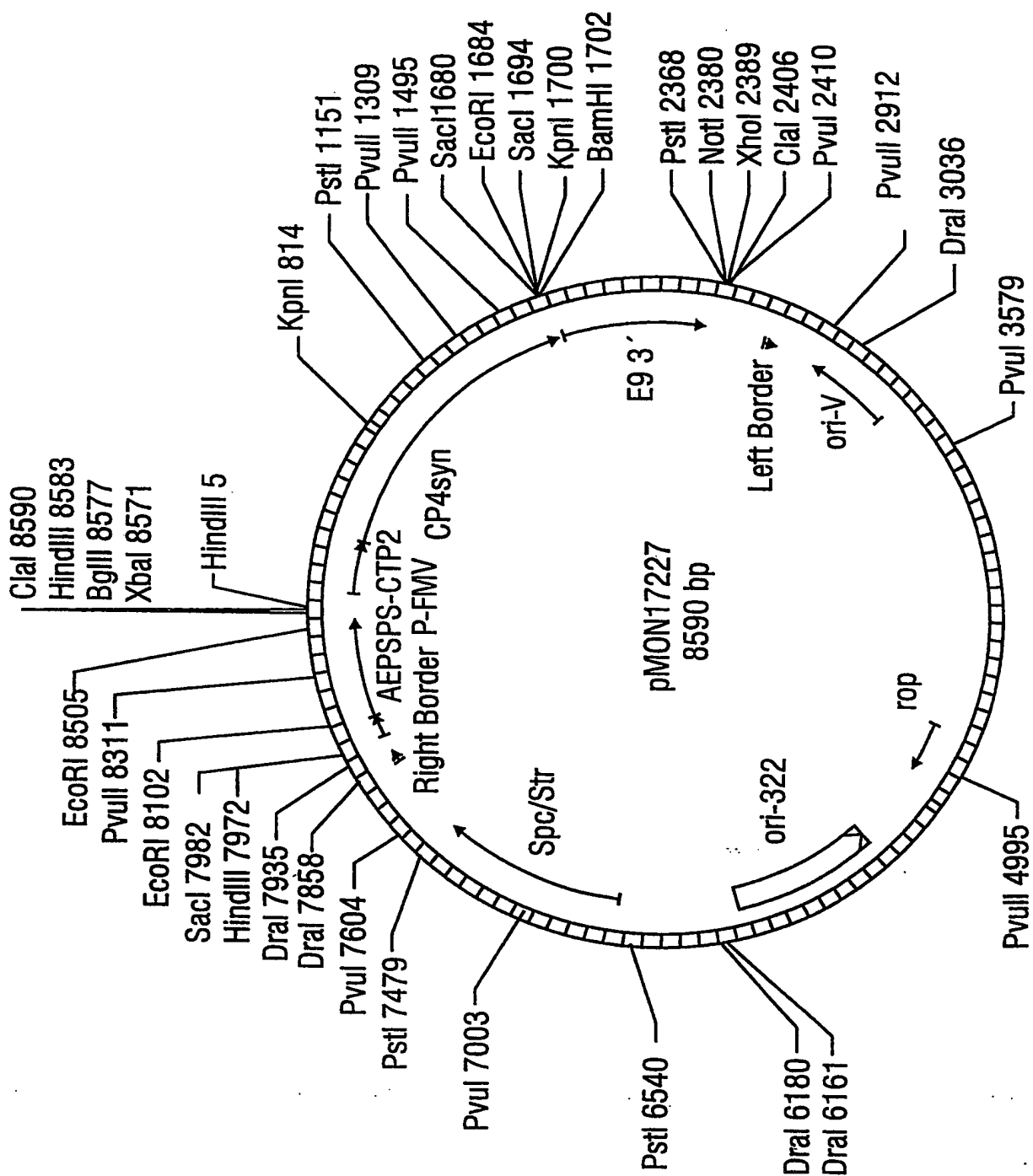
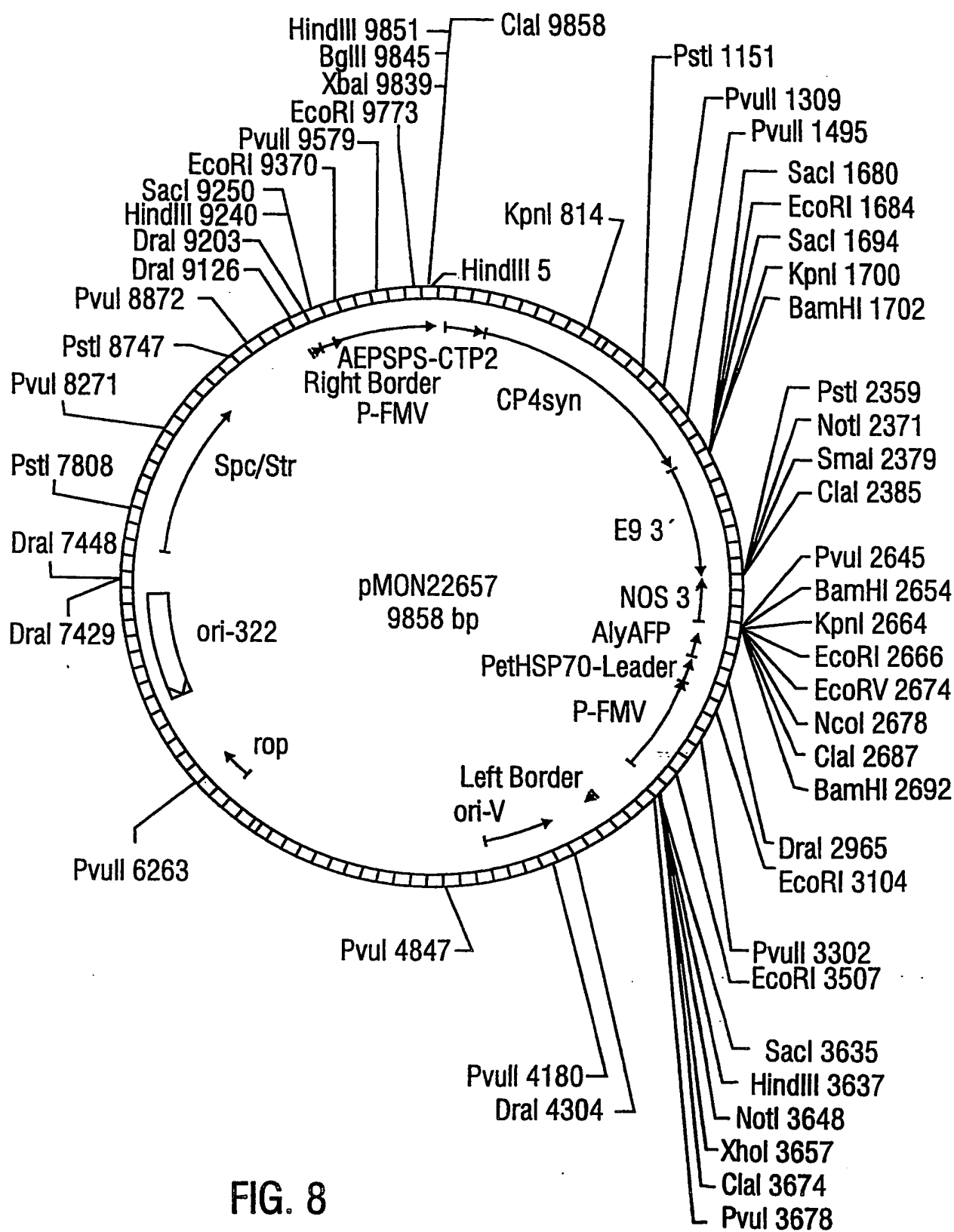
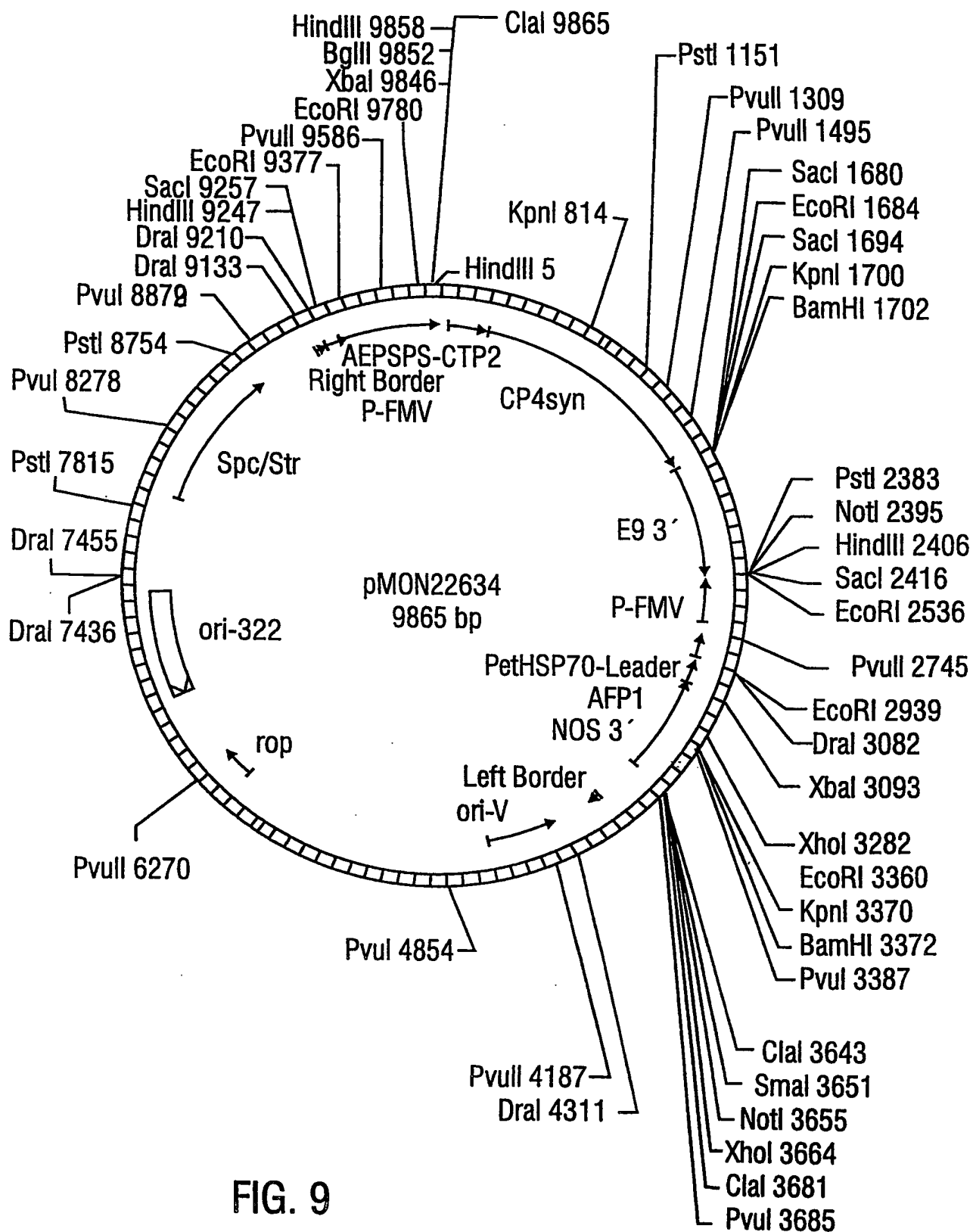


FIG. 7



**FIG. 8**





**FIG. 9**

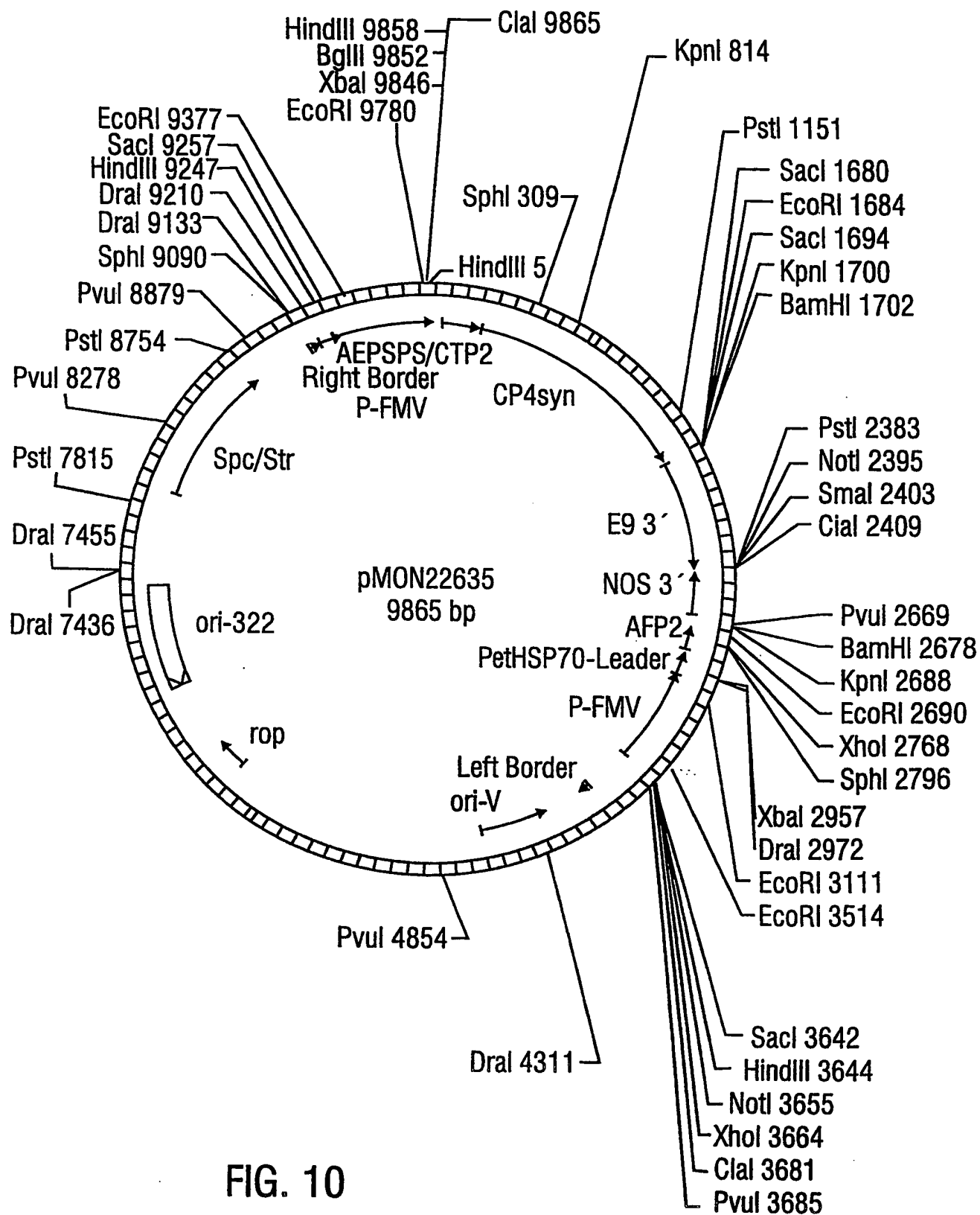


FIG. 10

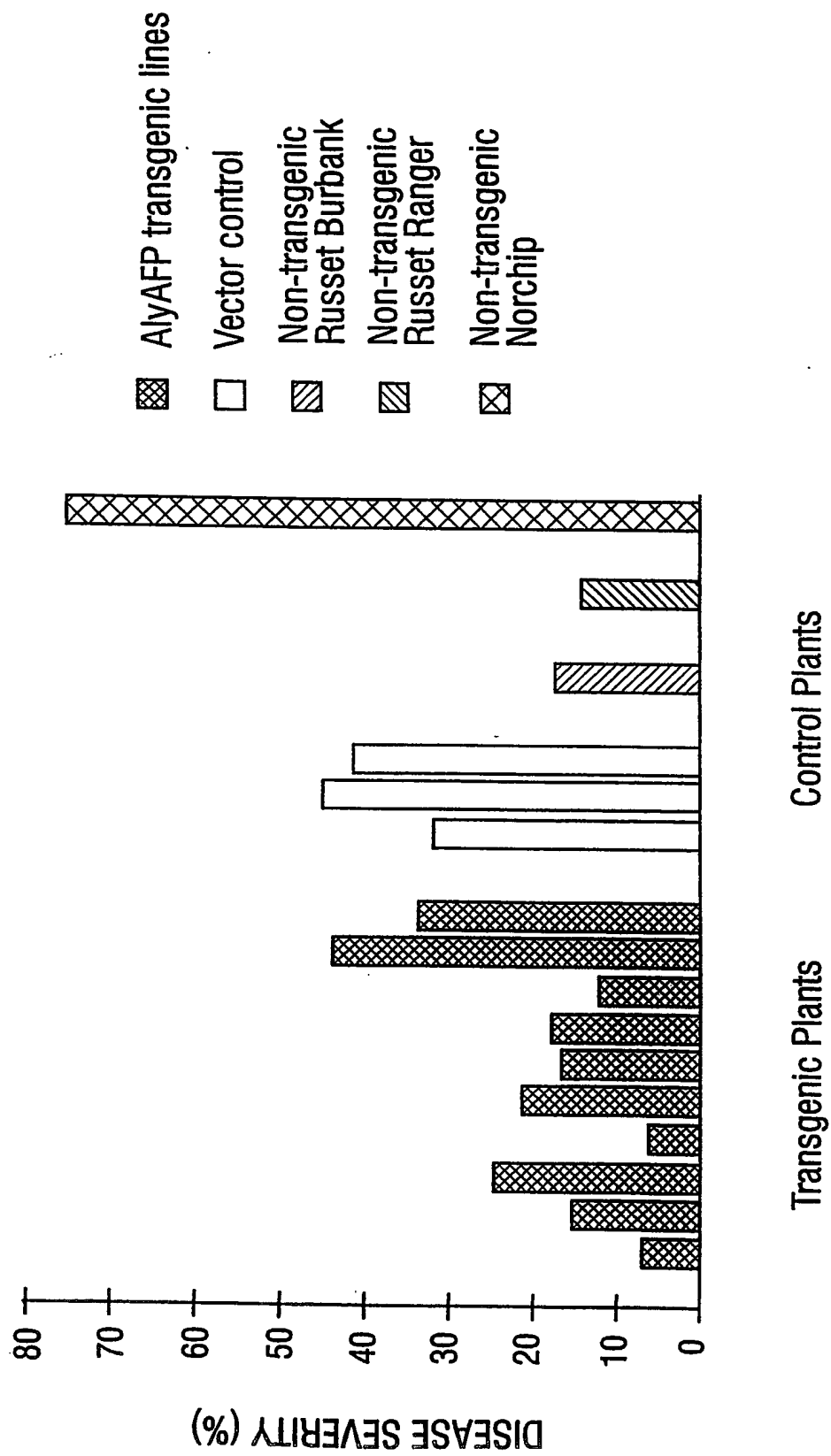


FIG.11

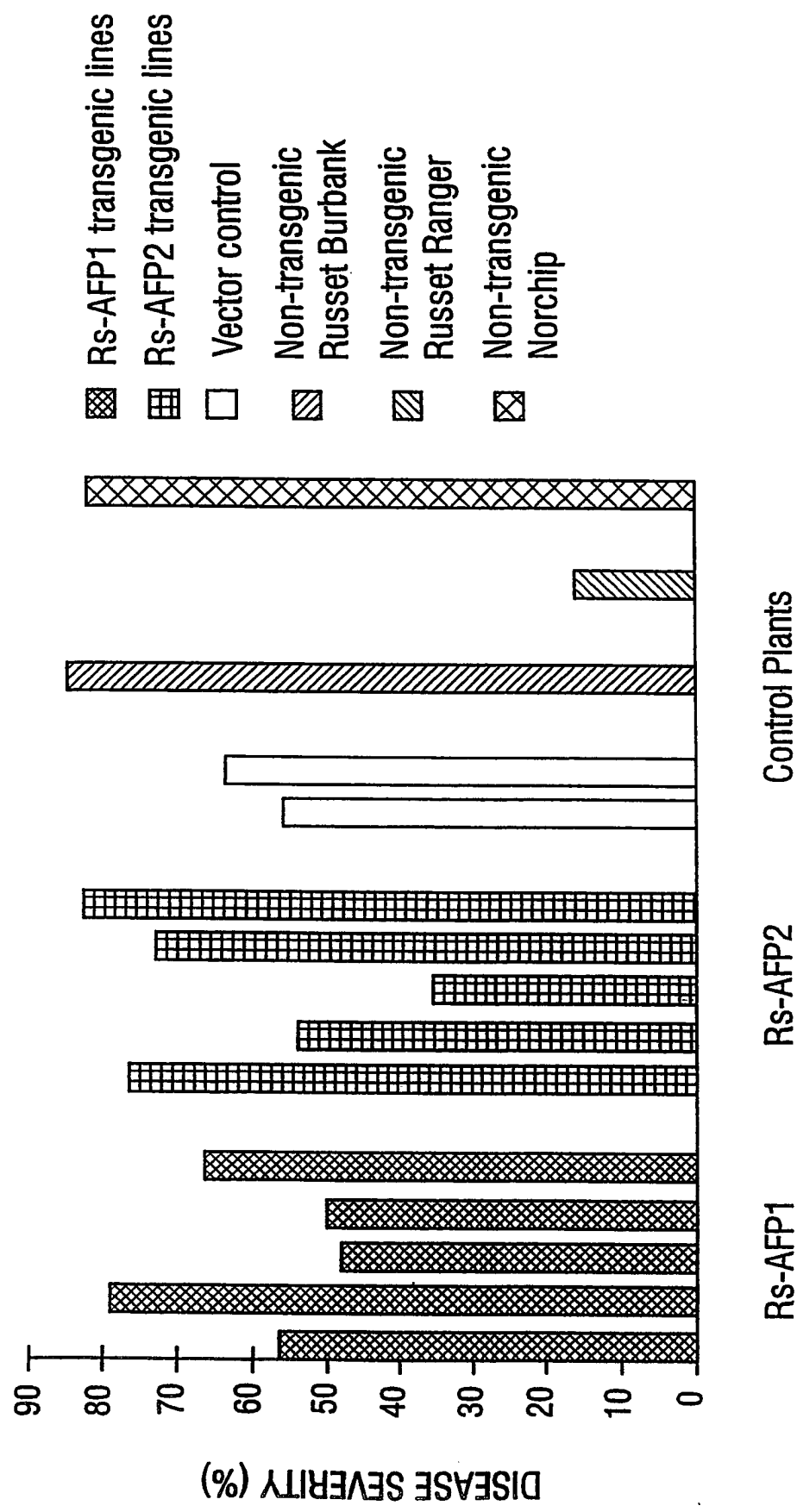


FIG.12